STUDENT ID NO										

MULTIMEDIA UNIVERSITY

FINAL EXAMINATION

TRIMESTER 2, 2017/2018

HPB 3021 - PARALLEL COMPUTING

(All Sections/Groups)

17 MARCH 2018 2:30 – 4.30 PM (2 Hours)

INSTRUCTIONS TO STUDENTS

- 1. This question paper consists of 3 pages, including this cover page.
- 2. You are required to attempt ALL four (4) questions. All questions carry equal marks (10).
- 3. Write all your answers in the Answer Booklet provided.

QUESTION 1 [10 MARKS]

a) List two features for the high performance computing.

[1 mark]

b) What is bioinformatics Linux distro? List down two bioinformatics linux distro.

[2 marks]

c) Write a bash shell script to find the biggest number from three given numbers. Numbers are read from the keyboard and the biggest number is displayed. [4 marks]

d) What is the function of X windows system?

[1 mark]

e) What is the advantage of top command for a multiuser environment?

[1 mark]

f) Describe the alternative command for "&".

[1 mark]

QUESTION 2 [10 MARKS]

a) A linux user observed that BLAST is not installed in the system. He applied apt-get install command but the error message prompted as shown in the figure below. Suggest two methods to solve the issue. [2 marks]

```
fls-mmu@flsmmu-OptiPlex-7040:~$ apt-get install mafft
E: Could not open lock file /var/lib/dpkg/lock - open (13: Permission denied)
E: Unable to lock the administration directory (/var/lib/dpkg/), are you root?
fls-mmu@flsmmu-CptiPlex-7040: -$ []
```

- b) Given a fasta formatted genome file (text file) namely "homo_sapien.fa", use Linux command to show the number of nucleotide adenine (A) in the file and the number of fasta sequences (>). [2 marks]
- c) What is the user from a group can do with 750 file permission?

[1 mark]

- d) Write the commands to download all the genome files from NCBI ftp and list out the downloaded file according to file sizes in ascending order . [3 marks] Assuming ftp link : ftp://ftp.ncbi.nlm.nih.gov/genomes/
- e) Write a shell script to run multiple sequence alignments for 100 sets of data.

[2 marks]

Continued...

QUESTION 3 [10 MARKS]

- a) Directory-based protocol is one of the approaches used to handle the cache coherence problem in distributed multiprocessors. Briefly explain directory protocols, with a suitable block diagram. [5 marks]
- b) Draw shuffle exchange networks with two, four and eight nodes. Make sure that you label the nodes. Is a shuffle exchange network with n nodes the same as a subgraph of a shuffle exchange network with 2n nodes? [5 marks]

QUESTION 4 [10 MARKS]

- a.) Write a complete C program using MPI library. The program initiates ranks and size of the processors and then every processor announces its rank (e.g. prints "This is the processor 1 of 10 processors"). [4 marks]
- b.) Briefly explain the following C programming commands in MPI library. Give a short description about the parameters of these functions.
 - i. int MPI_Send(void *buf, int count, MPI_Datatype dtype, int dest, int tag, MPI_Comm comm); lpt
 - ii. int MPI_Recv(void *buf, int count, MPI_Datatype dtype, int source, int tag, MPI_Comm comm, MPI_Status *status); 1pt
 - iii. MPI Abort (comm,errorcode); 1pt
 - iv. MPI_Probe (source,tag,comm,&status) 1pt

[4 marks]

c) What is the difference between "Asymmetric Multi-computers" and "Symmetric Multi-computers"?

[2 marks]

End of paper.

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